

Figure S1. cRNAsp12 predictions with the 16-26 bp dsRNAs highlighted by the dashed rectangles.

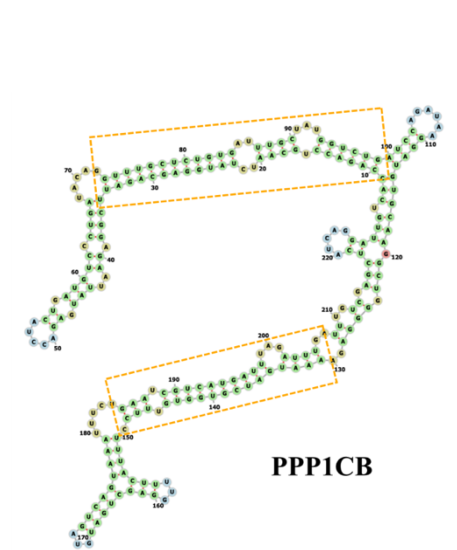
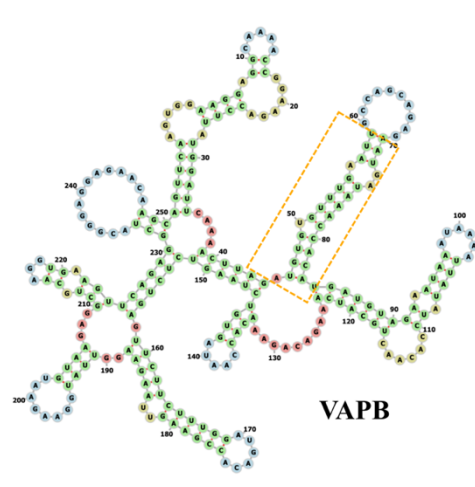
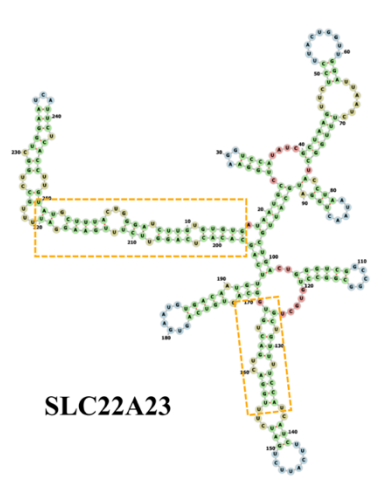
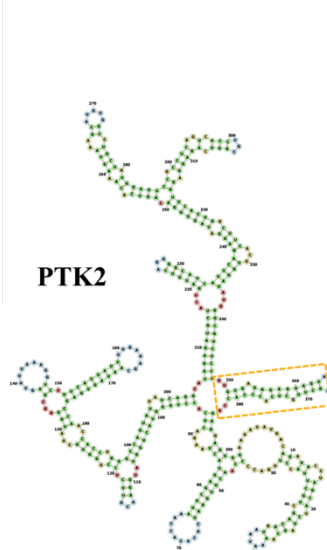
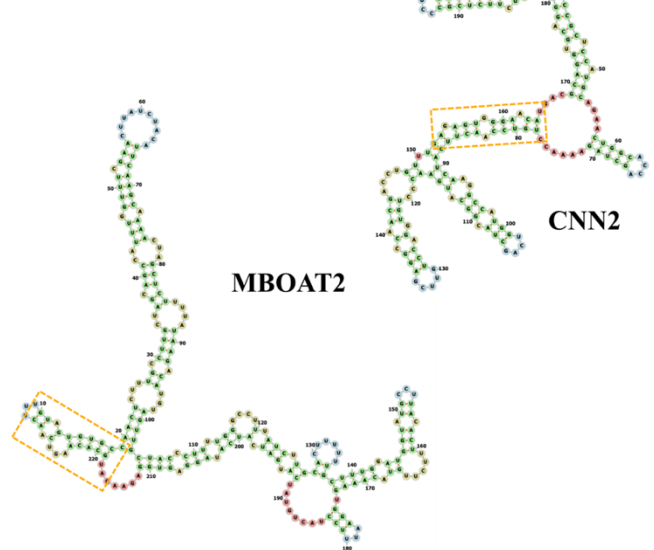
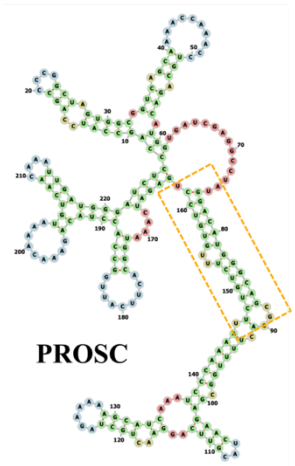
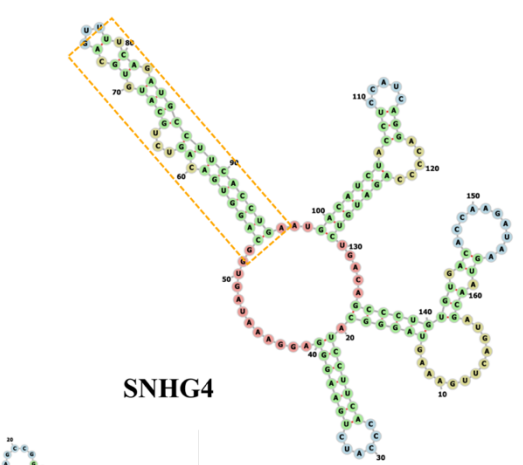
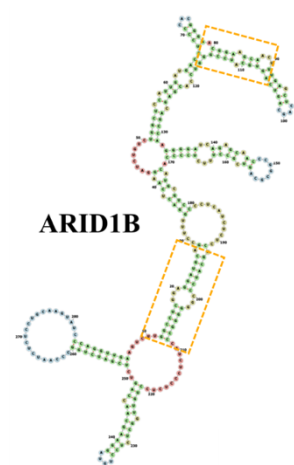
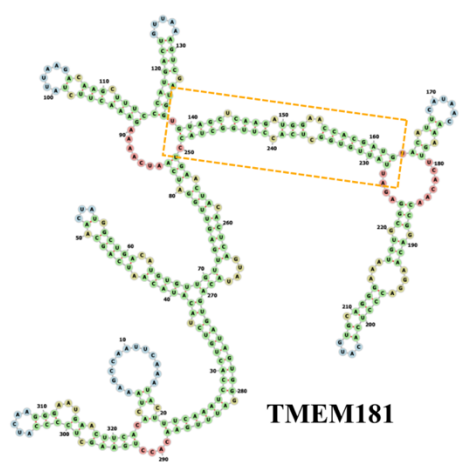


Figure S1. Continued.

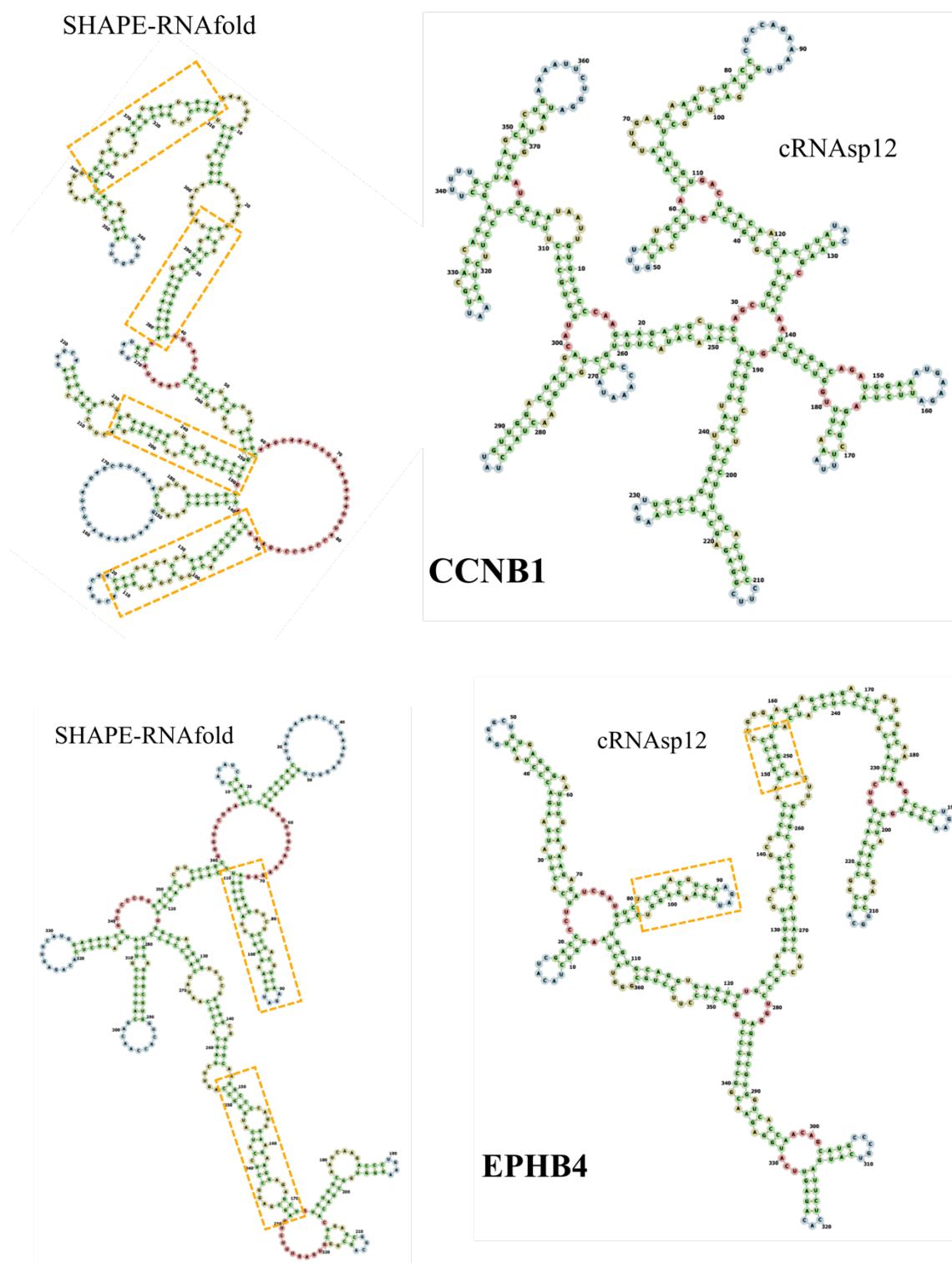


Figure S2. SHAPE-RNAfold and cRNAsp12 predictions with the 16-26 bp dsRNAs highlighted by the dashed rectangles.

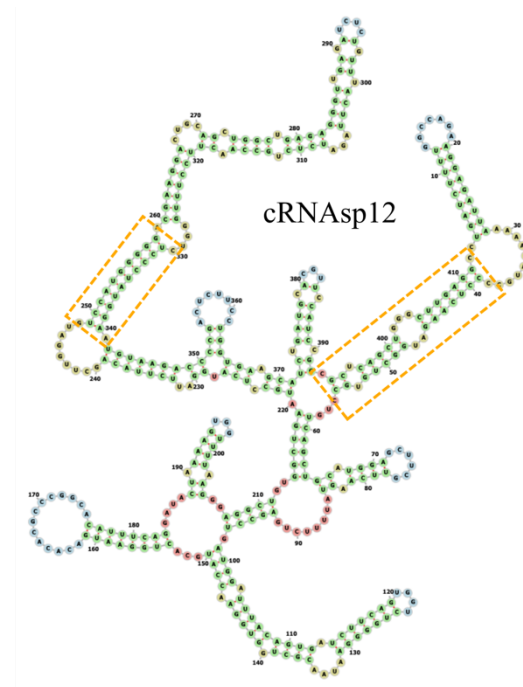
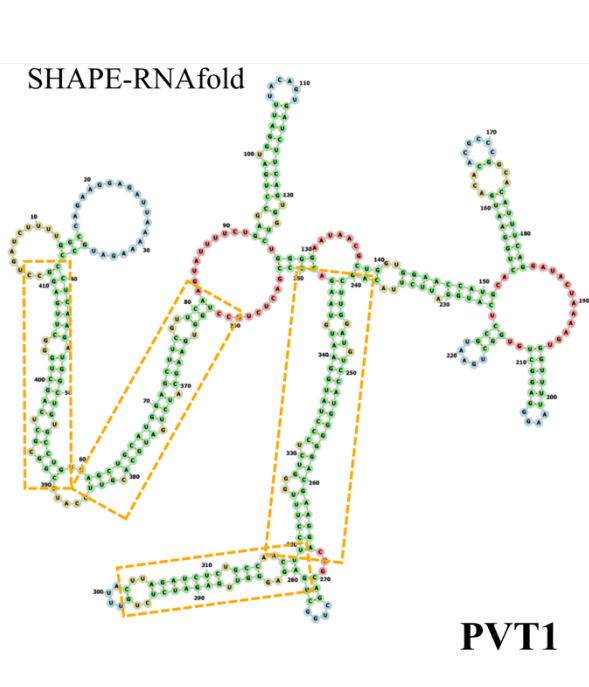
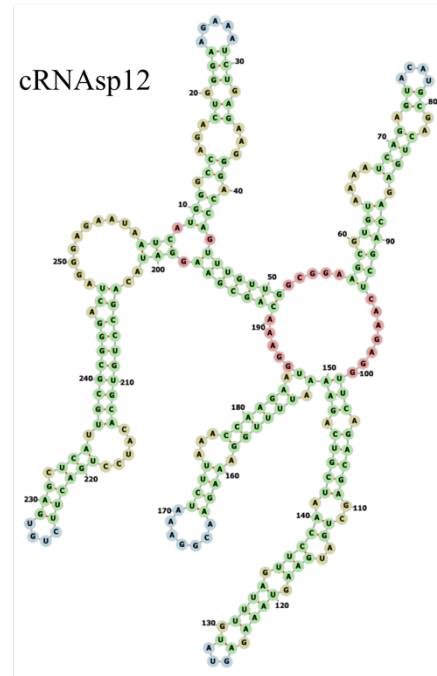
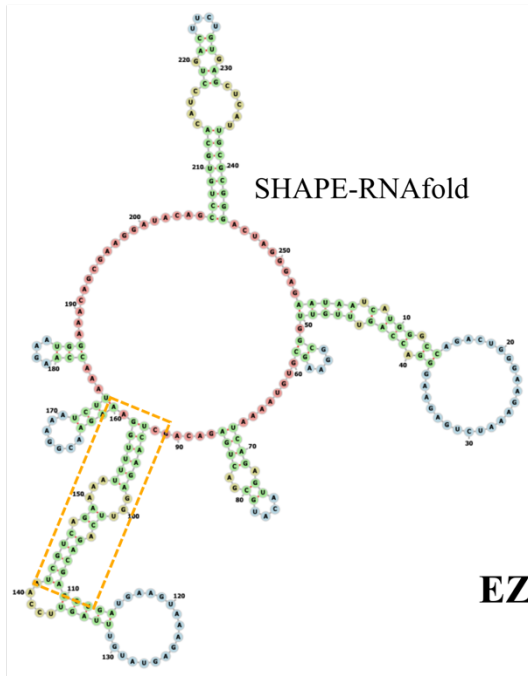


Figure S2. Continued.

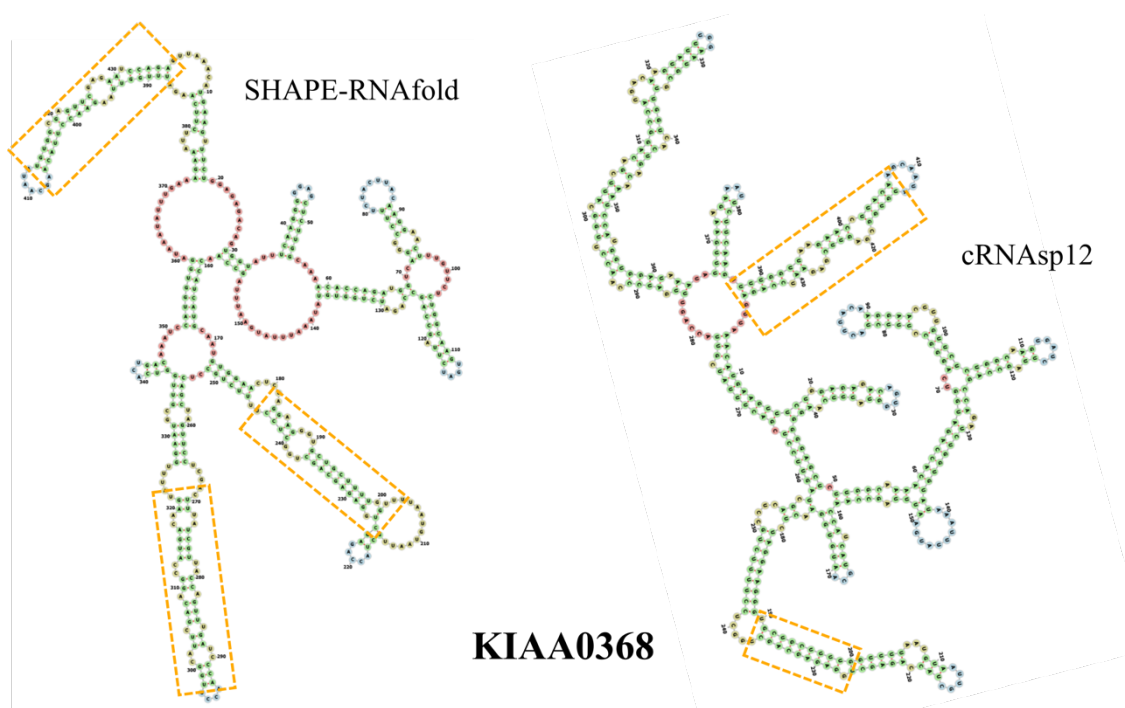
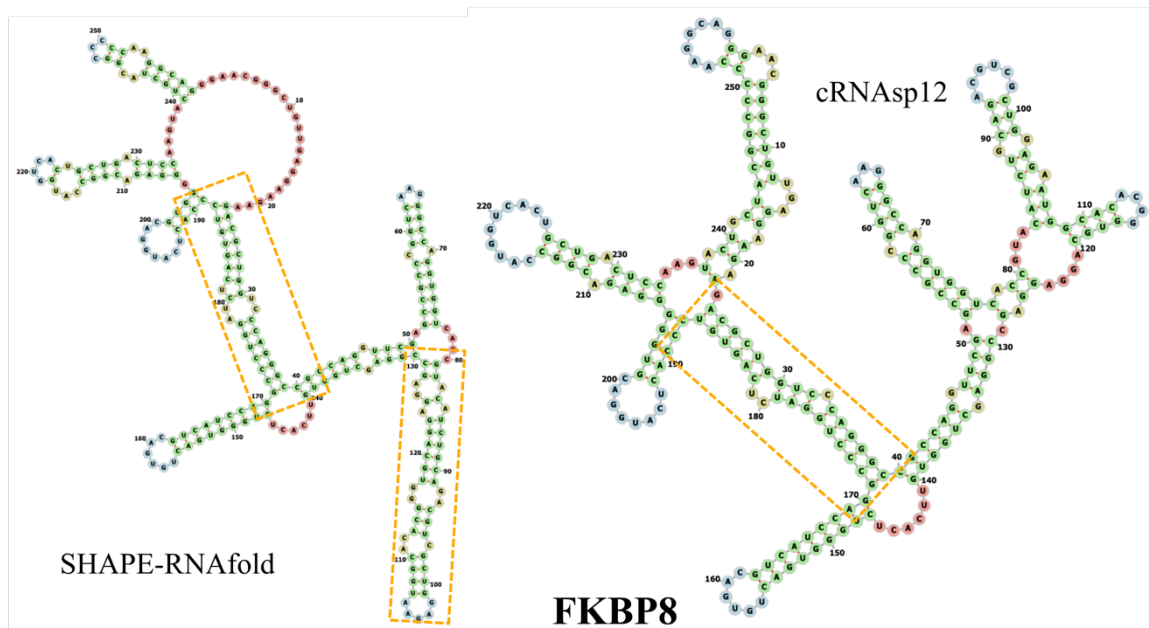


Figure S2. Continued.